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Sequence Listing was accepted.

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Reviewer: Anne Corrigan

Timestamp: [year=2008; month=2; day=27; hr=15; min=31; sec=21; ms=575;]
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Application No: 10537864 Version No: 2.1

Input Set:

Output Set:

Started: 2008-02-27 15:28:49.377
Finished: 2008-02-27 15:28:51.015
Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 638 ms
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Total Errors: 0
No. of SeqIDs Defined: 17
Actual SeqID Count: 17

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<110>   Mattson, Jeanine
        McClanahan, Terrill

<120>   Canine RANKL and Methods for Preparing and
        Using the Same

<130>   AH01646K

<140>   10537864
<141>   2005-06-07

<150>   US60/432092
<151>   2002-12-10

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<170>   PatentIn version 3.4

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ttcagggctc agatggatcc taatagaata tcagaagatg acactcactg cattaataga   180

attttcaaac tccatgaaaa tgcagatttg caagacacaa ctctggagaa tcaagacaca   240

aaattaatac ctgattcgtg taagagcatt aagcaggcct tccgagccgc cgtacaaaag   300

gaattacaac atattgttag atcacaacac atcagagcag aaaaagctat gatggaaggt   360

tcatggttgg aaatggccag gaggggcaag actcatactc aacctttgc tcatctcact   420

atcaatgcca ctgacatccc atctggttcc cacaaagtga gtctgtcctc ctggtaccat   480

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gaccgaggtt gggccaagat ctccaacatg actttcagca atgggaaact aatagttaac      540

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gacctcgcca cagagtatct tcagctgatg gtgtatgtca ctaaaaccag catcaaaatc      660

ccgagttctc atacactgat gaaaggaggt agcaccaaat actggtcagg gaattctgaa      720

ttccattttt attccataaa cgttggagga tttttaagc tacgatctgg tgaggaaata        780

agcatcgagg tatccaaccc atcactactg gacccagatc aagatgcaac atactttggg      840

gcttttaagg ttctagatat agattgagtc ccattttatg gagtgttatt ctgtatttcc      900

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ctaaggggta tgacccacaa tgatacaag                                        989



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Asp Pro Asn Arg Ile Ser Glu Asp Asp Thr His Cys Ile Asn Arg Ile
        35                  40                  45


Phe Lys Leu His Glu Asn Ala Asp Leu Gln Asp Thr Thr Leu Glu Asn
    50                  55                  60


Gln Asp Thr Lys Leu Ile Pro Asp Ser Cys Lys Ser Ile Lys Gln Ala
65                  70                  75                  80


Phe Arg Ala Ala Val Gln Lys Glu Leu Gln His Ile Val Arg Ser Gln
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His Ile Arg Ala Glu Lys Ala Met Met Glu Gly Ser Trp Leu Glu Met
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Ala Arg Arg Gly Lys Thr His Thr Gln Pro Phe Ala His Leu Thr Ile
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Trp Tyr His Asp Arg Gly Trp Ala Lys Ile Ser Asn Met Thr Phe Ser
145   150   155   160

Asn Gly Lys Leu Ile Val Asn Gln Asp Gly Phe Tyr Phe Leu Tyr Ala
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Asn Ile Cys Phe Arg His His Glu Thr Ser Gly Asp Leu Ala Thr Glu
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Tyr Leu Gln Leu Met Val Tyr Val Thr Lys Thr Ser Ile Lys Ile Pro
        195   200   205

Ser Ser His Thr Leu Met Lys Gly Gly Ser Thr Lys Tyr Trp Ser Gly
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Asn Ser Glu Phe His Phe Tyr Ser Ile Asn Val Gly Gly Phe Phe Lys
225   230   235   240

Leu Arg Ser Gly Glu Glu Ile Ser Ile Glu Val Ser Asn Pro Ser Leu
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Asp Ile Asp
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